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FORM PTO-1449 (REV. 7-85) U.S. DEPARTMENT OF COMMERCE PATENT AND TRADEMARK OFFICE

INFORMATION DISCLOSURE CITATION

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ATTY. DOCKET NO.
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APPLICATION NO.
10/509,009
APPLICANT
CHIQUET-EHRISMANN ET AL.
FILING DATE
SEPTEMBER 24, 2004

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EXAMINER INITIAL		DOCUMENT NUMBER	DATE	NAME	CLASS	SUBCLASS	FILING DATE
AG	AA	AA 6,124,260	Sharifi, et al.				
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FOREIGN PATENT DOCUMENTS

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AG	AM	WO 9421293		PCT				
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Sheet 2

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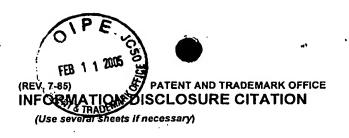
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· .	DE	Raouf, et al., "Discovery of Osteoblast-associated Genes Using cDNA Microarrays", Bone, Vol. 30 (2002)					
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be discovered, in one instance as a myotendinous antigen (Chiquet, M. & Fambrough, DM. (1984) J Cell Biol 98(6):1937-1946) and in another, as a protein enriched in the stroma of gliomas (Bourdon, MA. et al (1983) Cancer Res 43(6):2796-2805, reflecting the major sites of tenascin-C expression, namely in tendons and ligaments and the extracellular matrix of tumor stroma. A further instance of the discovery of tenascin-C (also termed hexabrachion) reflects its interaction with fibronectin (Erickson, HP. et al. (1984) Nature 311(5983):267-9). Enforced interaction of tumour cells with fibronectin can block proliferation in cell culture and can decrease tumour growth in nude mice (Akamatsu H. et al (1996) Cancer Res 56: 4541-4546 and Giancotti, F. G & Ruoslahti, E. (1990) Cell 60: 849-859). Tenascin-C was shown to disrupt the interaction of cells with fibronectin and in this manner may enhance tumour cell proliferation. Chiquet-Ehrismann, R. et al (1988) Cell <u>53</u>: 383-390 were the first to show that tenascin-C binds to fibronectin, blocks cell attachment to fibronectin and increases proliferation of rat breast adenocarcinoma cells (Chiquet-Ehrismann, R. et al (1996) Use 2011 131-139).

Tenascin-C is present in a large number of developments and tendons, it is absent from skeletal and heart muscle, unless the muscle has been injured. Tenascin-C expression is elevated in essentially all carcinomas as well as in many other types of tumors (for review see Chiquet-Ehrismann, R. (1993) Semin Cancer Biol 4(5):301-10). Furthermore, tenascin-C is upregulated in wound healing (Latijnhouwers, MA. et al. (1996) J Pathol 178(1):30-5), during skeletogenesis (Koyama, E. et al (1996) J Orthop Res. 14(3):403-412 and Hall, BK. & Miyake, T. (1995) Int J Dev Biol. 39(6):881-893) as well as in many diseases involving infections and inflammation (Schenk, S. et al. (1995) Int J Cancer 61(4):443-9).

30 Each tenascin family member exhibits a specific gene expression pattern during embryogenesis and in the adult (for review see Chiquet-Ehrismann, R. (1995) Experientia 51(9-10):853-62) suggesting specific roles for each member. Tenascin-R is an extracellular matrix component of the nervous system found mainly in brain tissue (Pasheva, P. et al. (2001) Prog Brain Res.

132:103-14. Review), whereas tenascin-X is prominently expressed in muscle and skin connective tissue. In one patient, tenascin-X deficiency has been reported to result in an Ehler's Danlos phenotype (Burch, GH. et al. (1997) Nat Genet 17(1):104-8).

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To date there is only one report on tenascin-W available in the literature. (Weber, P. et al. (1998) J Neurobiol 35(1):1-16). In this study, a cDNA encoding tenascin-W was isolated from a 20-28h postfertilization zebrafish cDNA library on the basis of the conserved epidermal growth factor-like domains found in all tenascin molecules. The expression pattern of tenascin-W transcripts was studied in the developing zebrafish by in situ hybridisation. It was found to be present in neural crest and sclerotome cells and the developing skeleton. Genebank sequence AJ001423 provides a zebrafish tenascin-W, and AL049689 provides a *novel human mRNA from chromosome 1, similar to Tenascin-R*, whose function is not known.

The present invention provides a compension comprising an isolated nucleic acid molecule having a nucleic science selected from the group consisting of:

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- (a) a nucleotide sequence as set forth in SEQ ID NO: 1;
- (b) a nucleotide sequence encoding the amino acid sequence shown in SEQ ID NO: 2;
- (c) a nucleotide sequence with at least 85% identity to the sequence of (a) or (b);

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- (d) a subsequence of more than 50 consecutive nucleotides of a sequence of (a), (b) or (c); and
- (e) a nucleotide sequence complementary to any of the nucleotide sequences or subsequence in (a),(b), (c) or (d).

In one aspect of the invention, the isolated nucleic acid molecule having a nucleotide sequence preferably exhibits at least 85% identity to the sequence of (a), more preferably encoding a variant of the amino acid sequence shown in SEQ ID NO: 2, such as a variant comprising an amino acid deletion, addition (e.g. fusion proteins) or substitution of the amino acid sequence